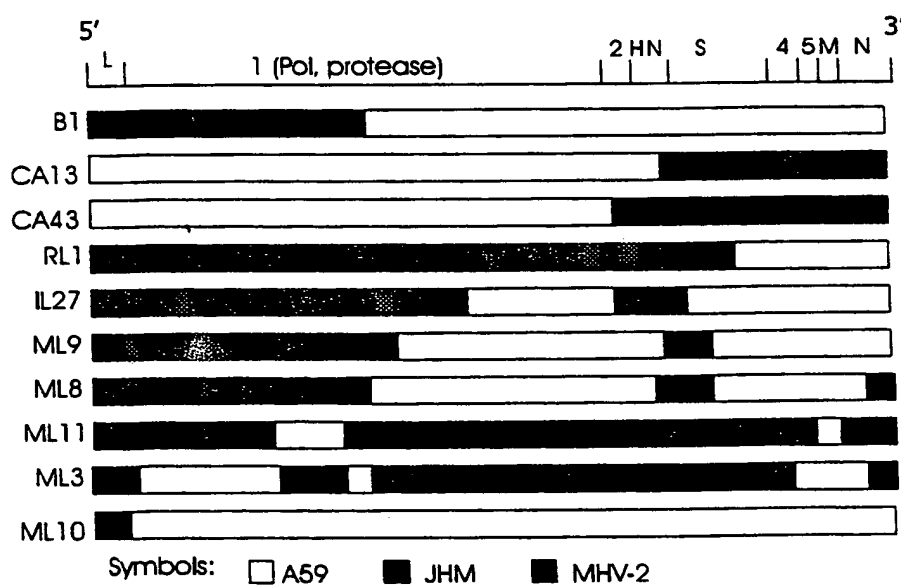
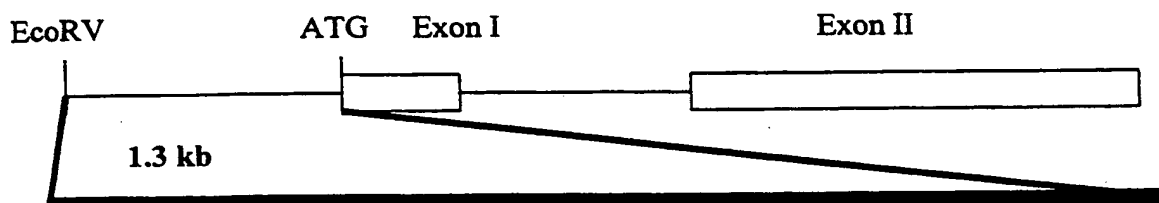


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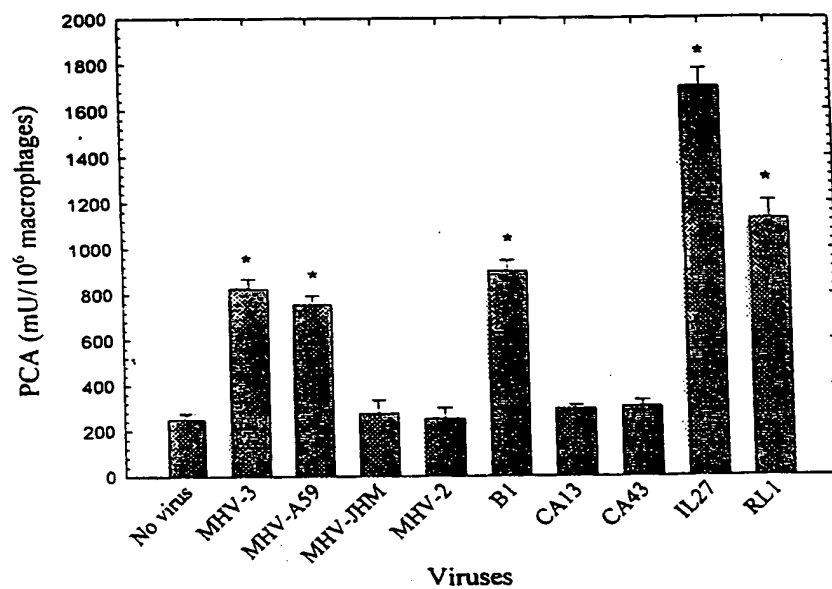
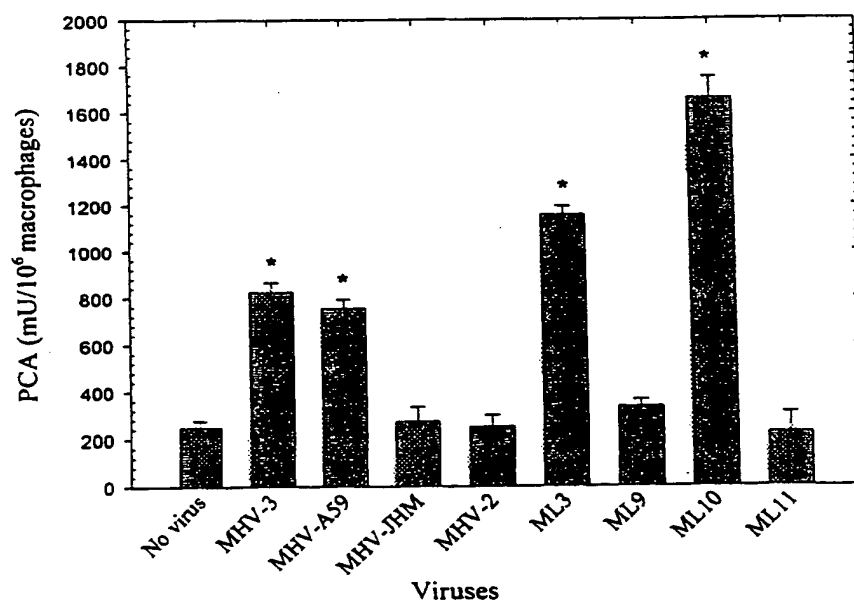
FIGURE 1

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FIGURE 2

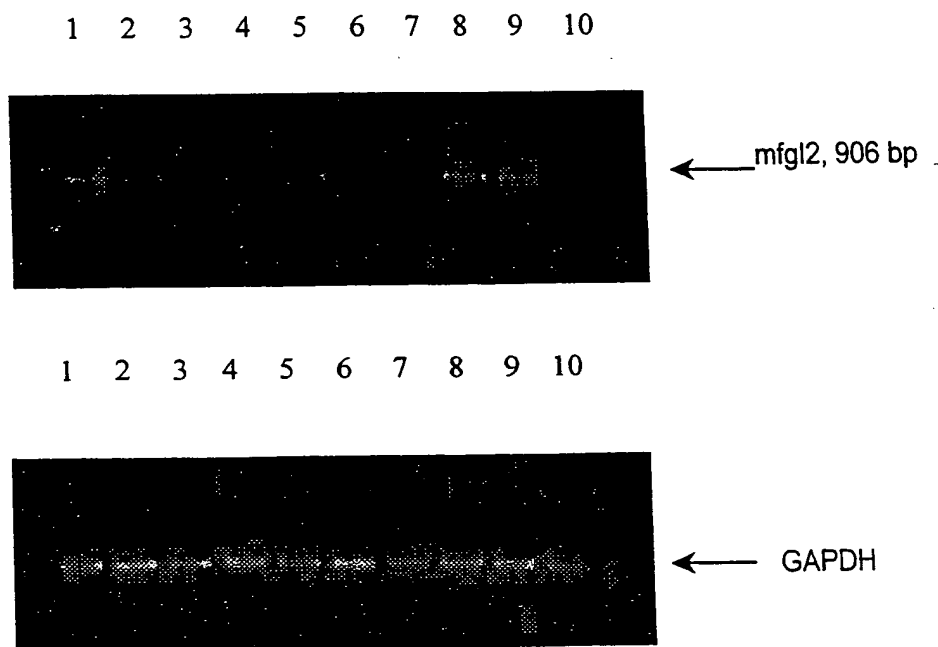
GATATCATGG	GATGGAATGA	GAAGGGAAAG	TAGGAGCCCG	AGAGTGCGGT	AAGACAAGGC	60
ATAAGGCGTG	TCTGACAAAT	TCTTCATACA	CACATTTCCC	CTTTGCACAT	TCAGTCTGTA	120
TAGGTTATTT	CTATAGGAGA	AAAAAATAT	TCAAATTCCT	TGTGCACTGG	TAACAGGCAT	180
GAAGGCTCAG	CAAAGCCAAT	ACGTGTTATG	TCCAGTTGGA	GACAGTGCCA	GGGCCAACAT	240
TCCAGACTTC	TCAGATAGAA	AGTGCGCCTG	CCTGCCCTGC	TCTGAGAATT	TGAAGAGAGT	300
AGTTCAGTTA	GAATTAAGAG	GCAGTAGAGA	AAAGTCTTGG	GAAATCTGGT	TAGAGATATA	360
AATATGAGAA	CTGGACATGG	TGGTACACAC	CTGTGATCTC	TGTGTTTAGG	AGGGAGAGGC	420
AGAGAGATCA	GGAGTTCAAG	GCCAGCCTGA	GCTACTTGAG	ACCCAGTCTA	AATAAATAAG	480
AGATAGATTA	CAGAGTGCCT	TTAACTAGTA	CAGAGAAAGA	ATTTGGGTTT	ATCTGTGTCA	540
GTTACGCTGA	AATAATTTTT	AAGTAATAAA	ATCCCTTTTA	ATAAGAAACC	TTATGAGGTC	600
AGTATGCACA	ATGAACTTAA	GAGAGACCCC	CAGTCTCTGA	GCTGAGTGAT	GGGGAAGGAC	660
AGCCACTGCC	TGTGATGTGT	GAGTGACGTG	CTTCCAAGTG	TTTTAACCAC	TGACGATTAC	720
ATAGCCTGCA	CAGTCAGGAG	AAAACAGCCG	TATTCTCTGC	CAGTTCTCTT	CCCTTTTACA	780
AACAGATGAG	AGACACACAC	AGAGAATCCA	TTTAAAGAGC	GGACCTTTGT	TCTGATTAGG	840
GGCAATTTTA	AGTACTTAAG	AGTTCACACA	AAGTCTAGCC	TTCAAAAAGA	AAACAGGTTC	900
CCAAACTAGG	GAGGAAACAG	AATCATTTCC	ATTTTGGTGA	CATTTAGTGG	GAAGAAGCTC	960
ACAGACATTT	AGACGTTCCA	ACTCTTTCCC	CACTAGTGGA	CCAAGTATAT	AATATGGTAT	1020
GMCSF		IE1.2	LF-A1			
CTTTTGGGCA	CTGGTATTAC	AACTGTTTTT	TAAACAAAAG	ACTTTCCTTG	TGCTTTACTA	1080
AAAACCCAGA	CGGTGAATCT	TGAATACAAT	GCGTGGCACC	CACGGCAGGC	ATTCTATTGT	1140
GCATAGTTTT	GACTGACAGG	AGATGACAGC	ATTTGGCTGG	CTGCGCTTGC	TGAGGACCCT	1200
CTCCTCCTGT	GTGGCGTCTG	AGACTGTGAT	GCAAATGCGC	CCGCCCTTTT	CTGGGAACTC	1260
		SP-1				
AGAACGCCTG	AGTCAGGCGG	CGGTGGCTAT	TAAAGCGCCT	GGTCAGGCTG	GGCTGCCGCA	1320
AP-1		TATA-box				
CTGCAAGGATG						

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FIGURE 3**A****B**

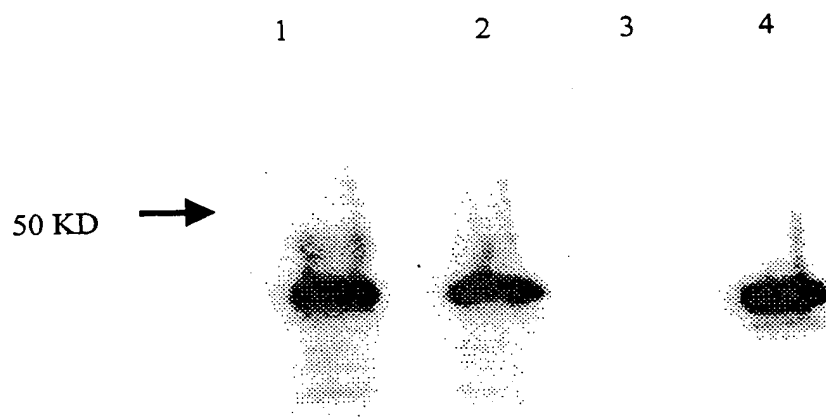
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FIGURE 4

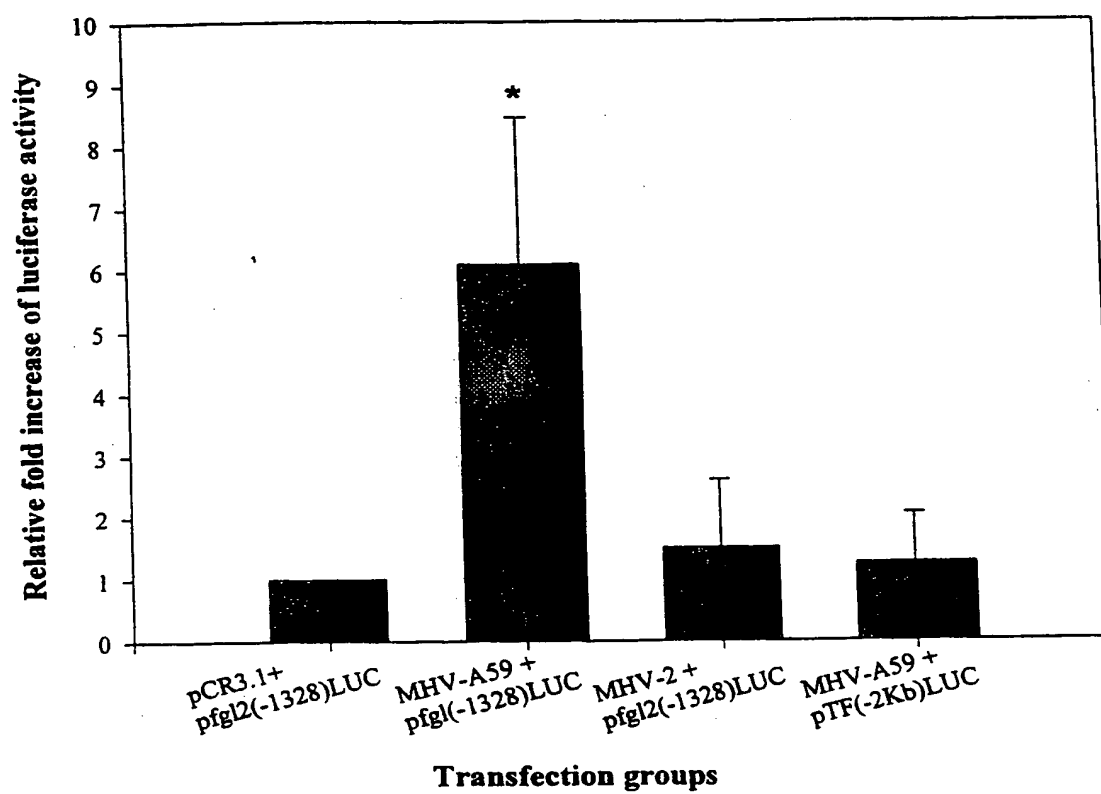


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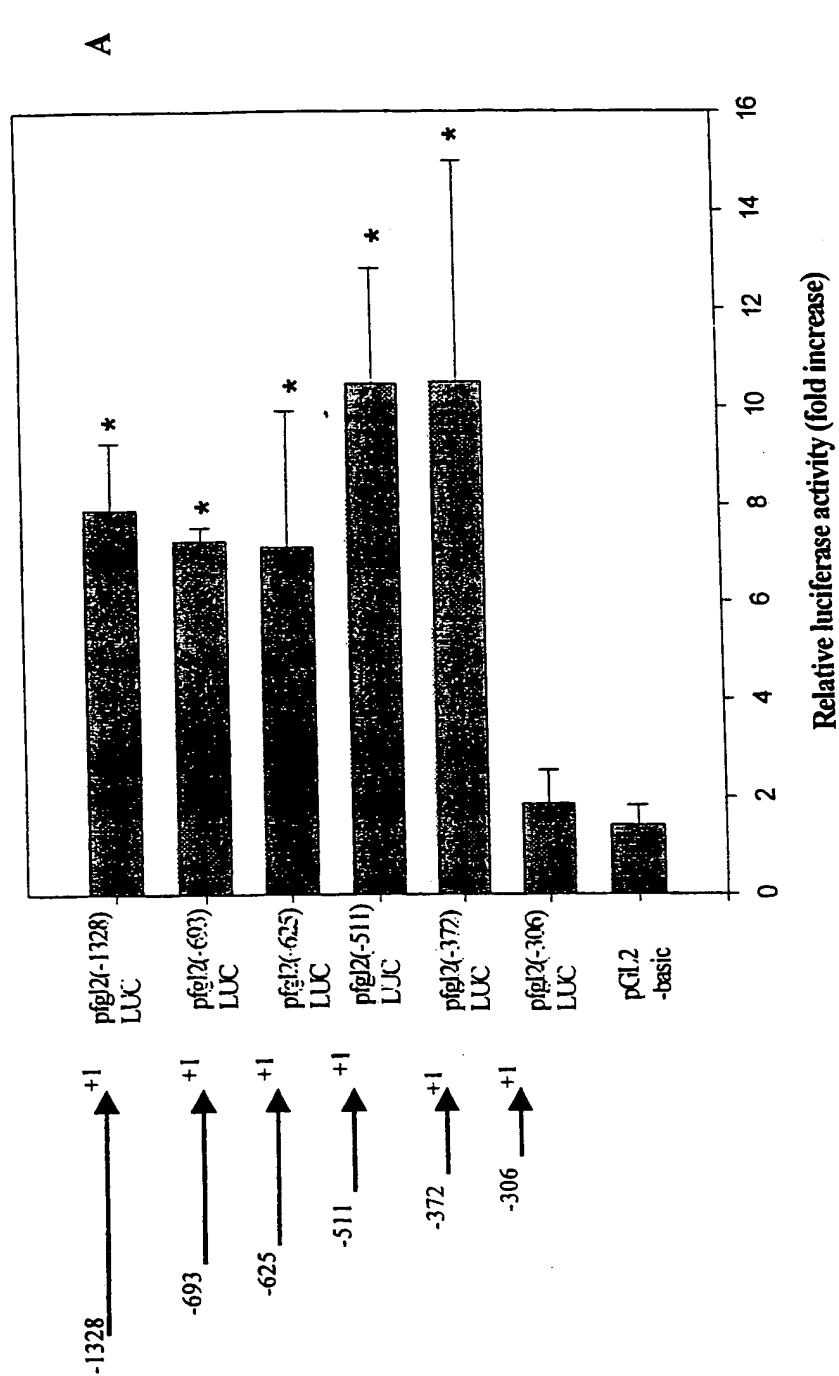
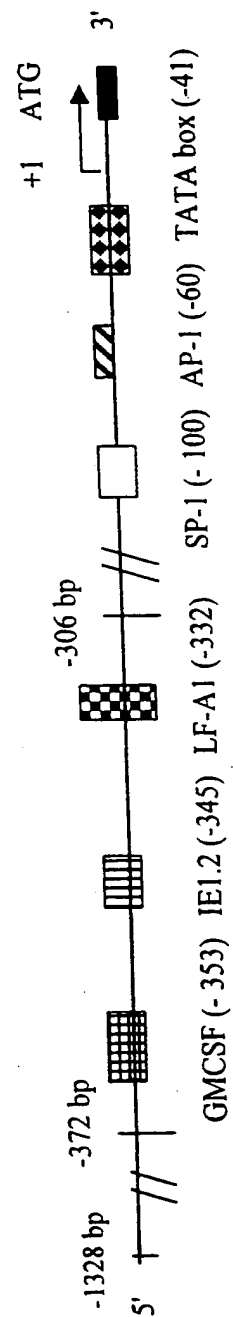
FIGURE 5



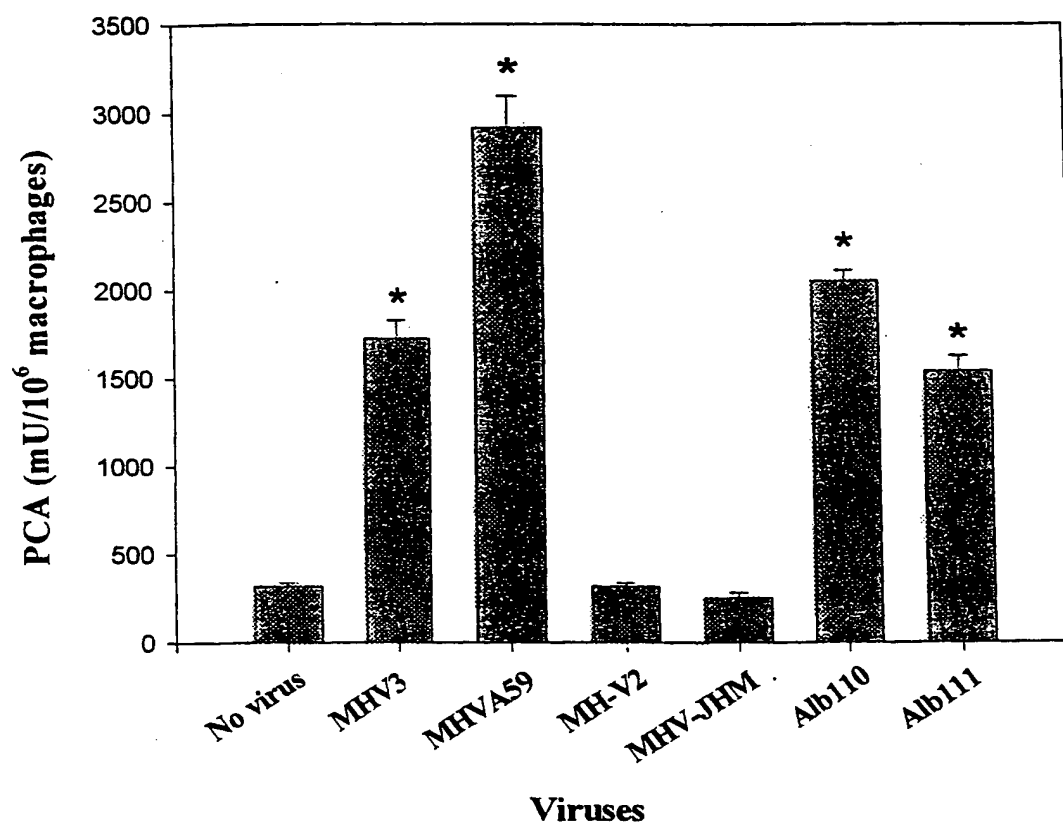
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FIGURE 6

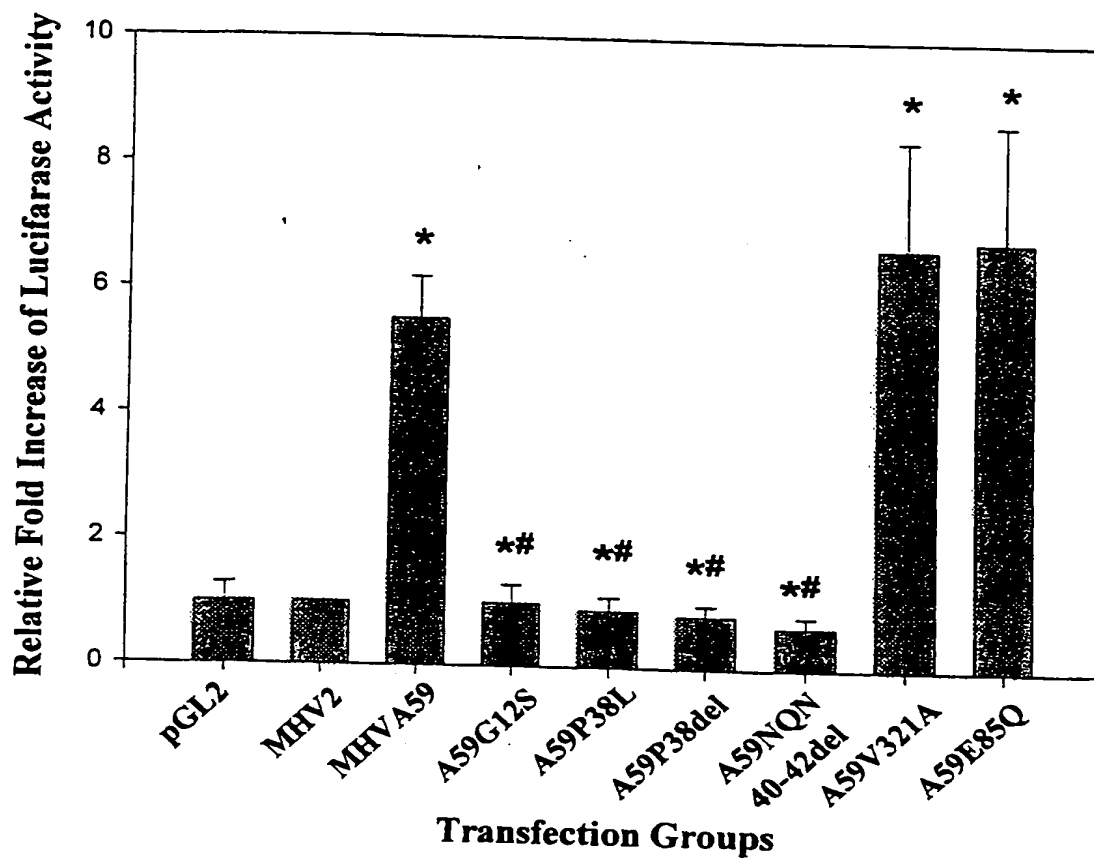
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FIGURE 7**B**

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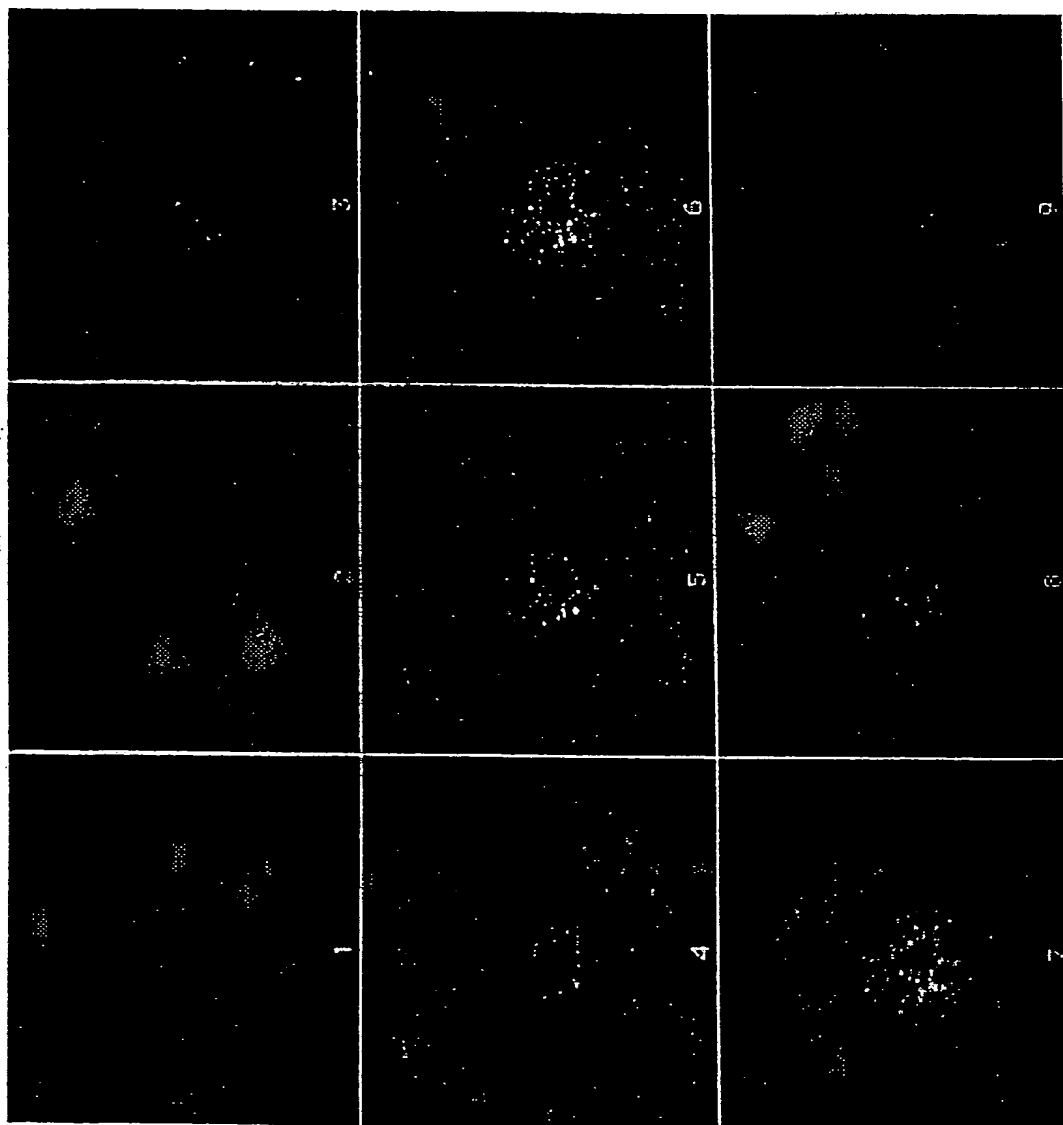
FIGURE 8

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FIGURE 9

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FIGURE 10



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FIGURE 11

1 2

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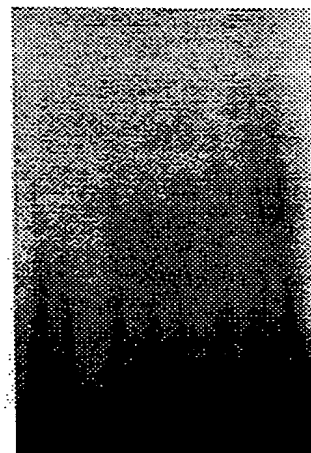
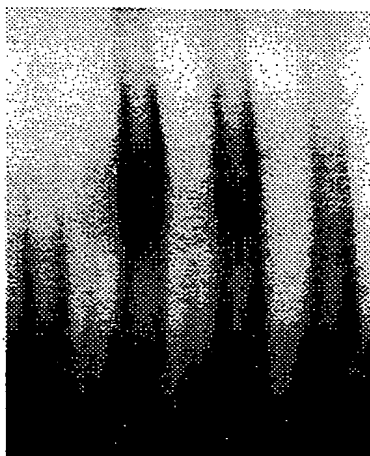
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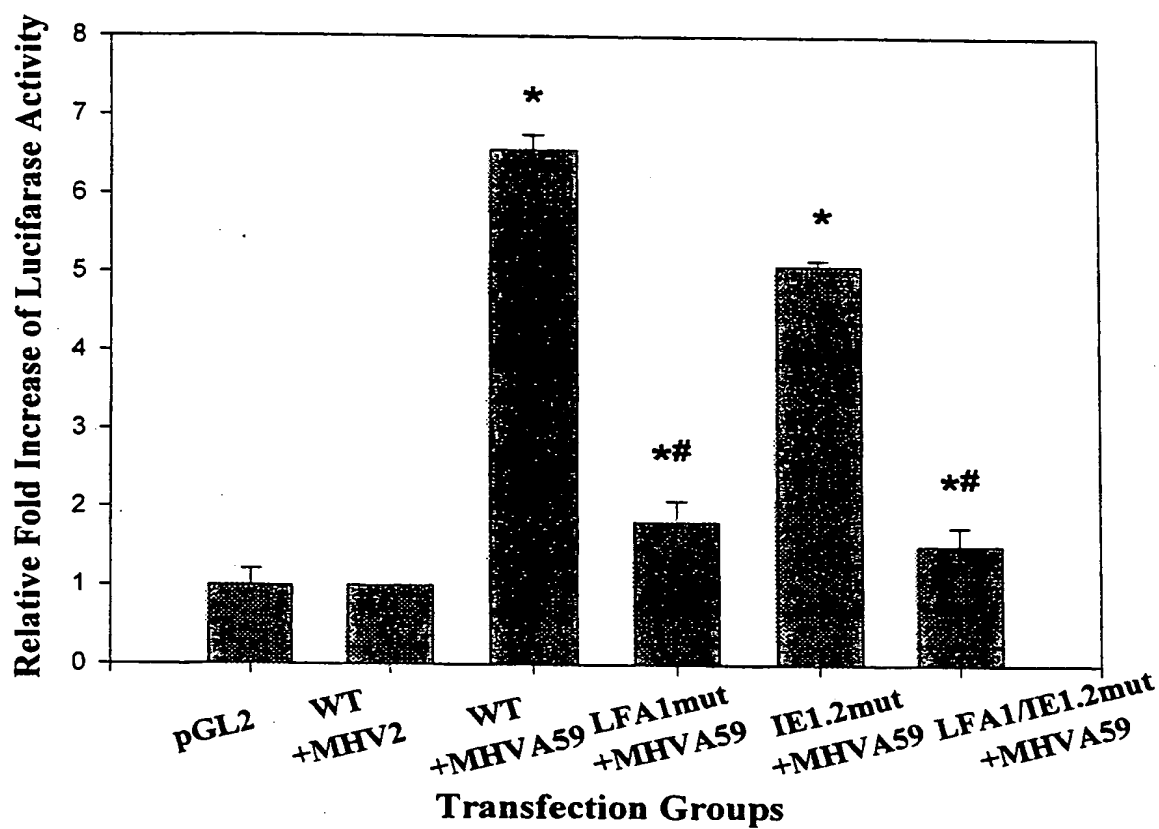
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FIGURE 12

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FIGURE 13

